

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/816,124

CRF Processing Date:

Edited by:

Verified by:

1600

1645

3/12/2002

 Changed a file from non-ASCII to ASCII Changed the margins in cases where the sequence text was "wrapped" down to the next line. Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____. Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as _____. Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____ Other:

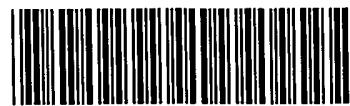
*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RECEIVED

TECH CENTER 1600/2900

MAR 15 2002



RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/816,124

DATE: 03/12/2002
TIME: 17:25:55

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\03122002\I816124.raw

3 <110> APPLICANT: Nagasawa, Yasuo
4 Yoshida, Hideaki
6 <120> TITLE OF INVENTION: Method for detecting and isolating genes
8 <130> FILE REFERENCE: S1-801PCT
10 <140> CURRENT APPLICATION NUMBER: US 09/816,124
11 <141> CURRENT FILING DATE: 2001-03-26
13 <150> PRIOR APPLICATION NUMBER: PCT/JP97/04126
14 <151> PRIOR FILING DATE: 1997-11-12
16 <150> PRIOR APPLICATION NUMBER: JP 1996-305163
17 <151> PRIOR FILING DATE: 1996-11-15
19 <160> NUMBER OF SEQ ID NOS: 4
21 <170> SOFTWARE: PatentIn version 3.0
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 29
25 <212> TYPE: DNA
26 <213> ORGANISM: Artificial Sequence
28 <220> FEATURE:
29 <223> OTHER INFORMATION: Artificially synthesized primer sequence
31 <400> SEQUENCE: 1 29
32 ataaagcttt cacatgagcg aaaaataaca
34 <210> SEQ ID NO: 2
35 <211> LENGTH: 29
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial Sequence
39 <220> FEATURE:
40 <223> OTHER INFORMATION: Artificially synthesized primer sequence
42 <400> SEQUENCE: 2 29
43 atggatccct attgtaaccc gcctgaagt
45 <210> SEQ ID NO: 3
46 <211> LENGTH: 38
47 <212> TYPE: DNA
48 <213> ORGANISM: Artificial Sequence
50 <220> FEATURE:
51 <223> OTHER INFORMATION: Artificially synthesized primer sequence
53 <400> SEQUENCE: 3 38
54 atgtctcgag aattcagtaa cccaggcatt attttatc
56 <210> SEQ ID NO: 4
57 <211> LENGTH: 30
58 <212> TYPE: DNA
59 <213> ORGANISM: Artificial Sequence
61 <220> FEATURE:
62 <223> OTHER INFORMATION: Artificially synthesized primer sequence
64 <400> SEQUENCE: 4

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/816,124

DATE: 03/12/2002
TIME: 17:25:55

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\03122002\I816124.raw

65 ttgtcctaga agcttgttg ctctgctgtc

30

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/816,124

DATE: 03/12/2002
TIME: 17:25:56

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\03122002\I816124.raw



1645

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/816,124

DATE: 03/01/2002
TIME: 15:07:13

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\03012002\I816124.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Nagasawa, Yasuo
4 Yoshida, Hideaki
6 <120> TITLE OF INVENTION: Method for detecting and isolating genes
8 <130> FILE REFERENCE: S1-801PCT
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13 <150> PRIOR APPLICATION NUMBER: PCT/JP97/04126
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17 <151> PRIOR FILING DATE: 1996-11-15
19 <160> NUMBER OF SEQ ID NOS: 4
21 <170> SOFTWARE: PatentIn version 3.0

ERRORED SEQUENCES

56 <210> SEQ ID NO: 4
57 <211> LENGTH: 30
58 <212> TYPE: DNA
59 <213> ORGANISM: Artificial Sequence
61 <220> FEATURE:
62 <223> OTHER INFORMATION: Artificially synthesized primer sequence
64 <400> SEQUENCE: 4
65 ttgtcctaga agcttgtgtc ctctgctgtc
E--> 67 38
E--> 69 1/2

30

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/816,124

DATE: 03/01/2002
TIME: 15:07:14

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\03012002\I816124.raw

L:67 M:254 E: No. of Bases conflict, LENGTH:Input:38 Counted:30 SEQ:4
M:254 Repeated in SeqNo=4
L:69 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:69 M:252 E: No. of Seq. differs, <211>LENGTH:Input:30 Found:31 SEQ:4